

FIGURE 1A

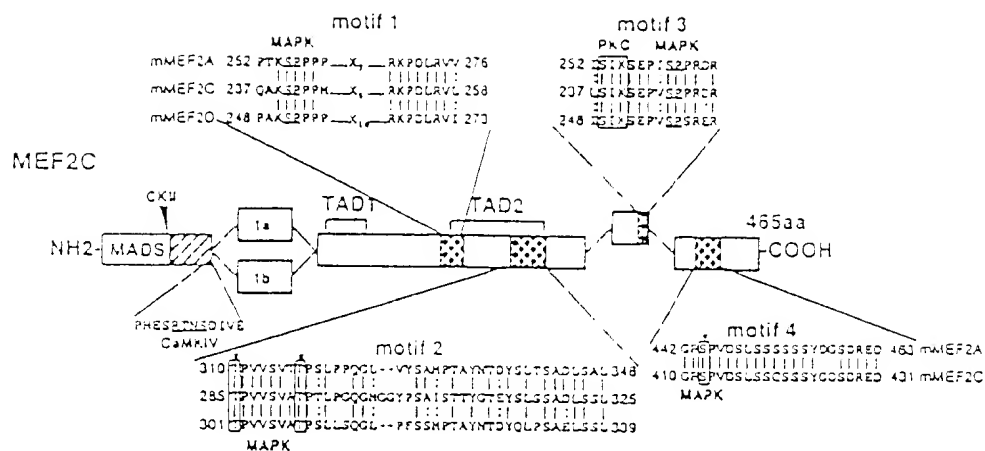


FIGURE 13

[illegible]

FIGURE 2A

MGRKKIQITRIMDERNRQVTFTKRKEGLMKKAYELSVLCCCEIALIIFNSSNKLFQYASTDM
DKVLLKYTEYNEPHESRTNSDIVEALNKKEHRGODSPCPOTSYVLTPTHTEKYKKINEEFDN
MMRNHKKIAPGLPPQNFMSMTVPVTSFNALSNTNPGSSLVSPSLAASSTLTSSMSPPPQT
LHRNVSPGAPQRRPPSTGNAGGMLSTTDLTVPNGAGSSFVGNGFVNSRASPNLIGATGANSIG
KVMPTKSPPPPGGGLGMNSRKPDLRVVIPTSSKGMMPPLSEEELELNTQRISSSQATQPL
ATPVVSVTTPSLPPQGLVYSAMPTAYNTDYSLTSAOLBALQGFSNPGMLSLGQVSAWQQHHL
GQAALSSLVAGGQLSQGSNLSINTNQNISIKSEPISPFRDRMTPSGFQQQQQQQQQQQPPPP
PQPQPQPQPQPQRQEMGRSPVDSLSSSSSSSYDGSOREDPRGDFHSPVILGRPPNTEDRES
PSVKRMRMDAWVT

FIGURE 2B

FIGURE 3A

MGRKKIQISRILDQNRQVTFTKPKFGLMKKAYELSVLCDCETALIIFNSANRLFQYASTDM
DRVLLKYTEYSEPHESTRNTDILETLKRRGIGLDGPELEPDEGPPEEPGEKFRRLAGEGGDPA
LPRPRLYPAAPAMPSPDVVYGALPPPCCDPSGLGEALPAQSRPSPFRPAAPKAGPPGLVHPL
FSPSHLTSKTPPPLYLPTEGRRSDLPGGLAGPRGGLNTSFSLYSGLQNPSTATPGPPLGSF
PFLPGGPPVGAELAWARRVPQPAAPRRPPQSASSLSASLPFGAPATFLRPSPIPCSSPGPW
QSLCGLGPPCAGCPWPTAGPGRRSPGGTSPERSPGTARAGDPTSLQASSEKTQQ

FIGURE 3B

FIGURE 4A

FIGURE 4A CONTINUED

MGRKKIQITRIMDERNRQVTFTKRKFGLMKKAYELSVLCDCEIALIIFNSTNKLQYASTOM
DKVLLKYTEXNEPHESRTNSDIVETLRKKGLNGCDSPPDPAODSVGHSPESDOKYRKINEDI
DLMISRQRLCAVPPPNFEMPVSI PVSSHNSLVYSNPVSSLGPNLLPLAHPSLQRNSMSPGV
THRPPSAGNTGGLMGGDLTSGAGTSAGNGYGNPRNSPGLLVSPGNLNKNMQAKSPPPMNLGM
NNRKPCLRVLIIPPGSKNTMPSVSEDDVLLINQRINNSQSAQSLATPVVSVATPTLPGQGMGG
YPSAISTTYGTEYSLSSADLSSLSGFNTASALHLGSVTGWQQQLHNMPPSALSQLGACTST
HLSQSSNLSLPSTQSLNIKSEPVSPPRDRITTPSRYPQHTRHEAGRSPVDSLSSCSSSYDGS
DREDHRNEFHSPIGLTRPSPDERESPSVKRMRLSEGWAT

FIGURE 4B

[illegible]

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[illegible]

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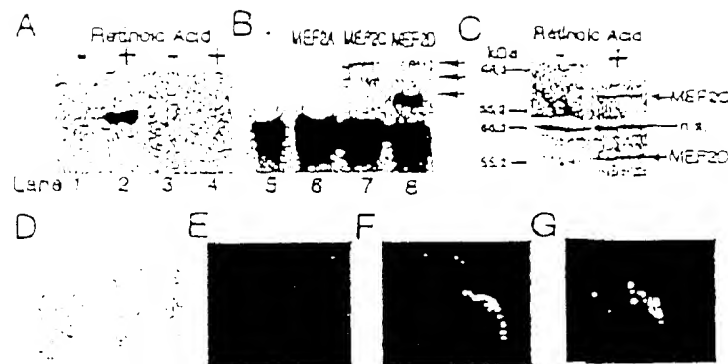


FIGURE 6

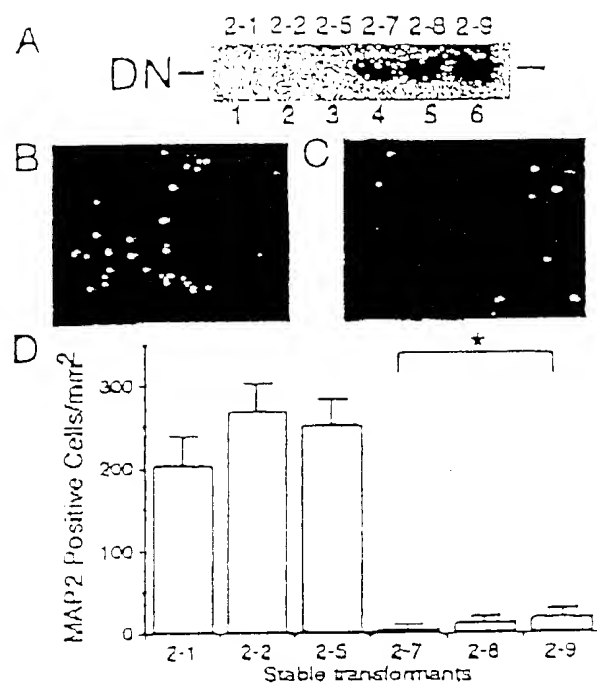


FIGURE 7

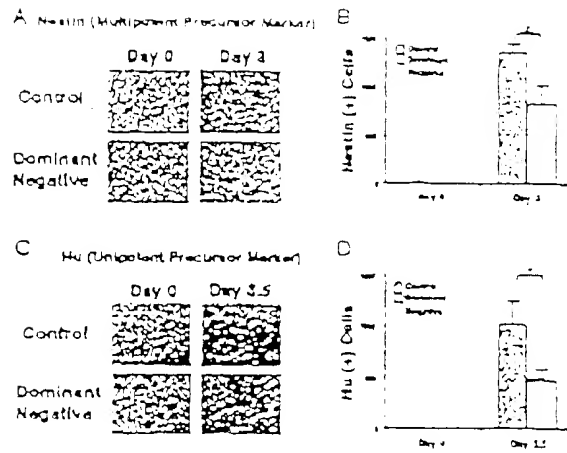


FIGURE 8

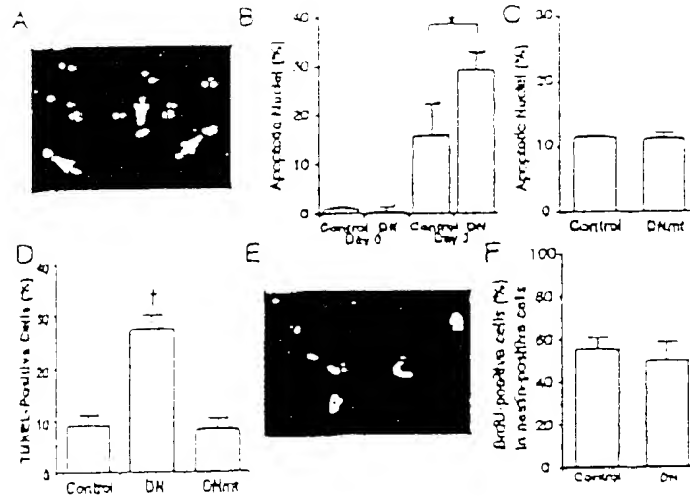


FIGURE 9

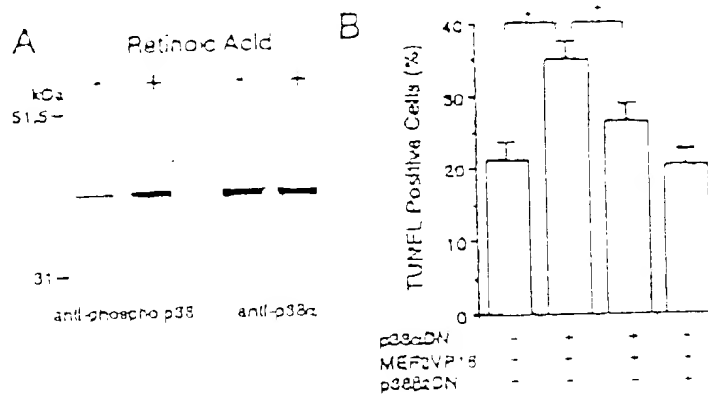


FIGURE 10